

Figure 1

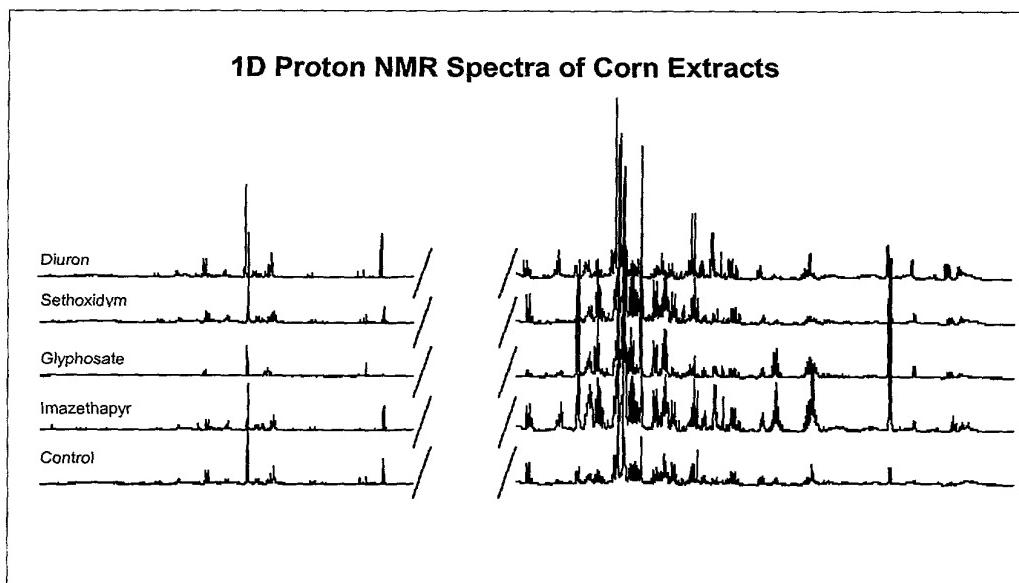


Figure 2

Batch_Spectrum	Seq. No.	Treatment	NetworkOutput: Node/Node Value	Network Output Assignment
Training Set				
na022400_02	2	Control	0.99996	0.00003
na022400_05	5	Control	0.99998	0.00002
na022400_06	29	Control	0.99998	0.00003
na030100_09	32	Control	0.99998	0.00002
na030100_11	34	PURSUIT	0.99996	0.00001
na030100_14	37	PURSUIT	0.99996	0.00004
na030100_17	40	PURSUIT	0.99995	0.00002
na030100_19	42	PURSUIT	0.99995	0.00004
na030600_04	51	Control	0.99993	0.00001
na030600_08	55	Sethoxydim	0.00002	0.00002
na030600_10	57	Sethoxydim	0.00002	0.00005
na030600_13	60	Foul	0	0.00002
na030600_15	62	Glyphosate	0.00007	0
na030600_16	63	Glyphosate	0.00006	0.00003
na030600_20	67	Diuron	0.00004	0.00003
na030600_21	68	Diuron	0.00007	0.00003
Test Set				
na022400_01	1	Control	0.99998	0
na022400_03	3	Control	0.99996	0
na022400_04	4	Control	0.99998	0
na022400_06	6	Control	0.99997	0
na022400_07	7	Control	0.99998	0
na022400_08	8	Control	0.99997	0
na022400_09	9	PURSUIT	0.02733	0
na022400_10	10	PURSUIT	0.00086	0.00011
na022400_11	11	PURSUIT	0.00085	0.00141
na022400_12	12	PURSUIT	0.00016	0.0025
na022400_13	13	PURSUIT	0.00013	0.00259
na022400_14	14	PURSUIT	0.00013	0.00238
na022400_15	15	PURSUIT	0	0.10229
na022400_16	16	PURSUIT	0.00005	0.00469
na022400_17	17	PURSUIT	0.00114	0.001127
na022400_18	18	PURSUIT	0.00091	0.00139
na022400_19	19	PURSUIT	0.00048	0.00185
na022400_20	20	PURSUIT	0.00214	0.00104
na022400_21	21	PURSUIT	0.00073	0.00088
na022400_22	22	PURSUIT	0.00046	0.00146
na022400_23	23	PURSUIT	0.00086	0.00123
ra030100_01	24	Control	0.99996	0
ra030100_02	25	Control	0.99997	0
ra030100_03	26	Control	0.99997	0
ra030100_04	27	Control	0.99996	0
ra030100_05	28	Control	0.99993	0

Figure 3a

Training Set		Control	PURSUIT	Sethoxydim	Glyphosate	Diuron	Foul	Assignment
na030100_07	30	Control	0.99998	0	0.00002	0.00002	0.00001	Control
na030100_08	31	Control	0.99996	0	0.00002	0	0.00001	Control
na030100_10	33	PURSUIT	0	0.99996	0.00001	0.00003	0.00005	PURSUIT
na030100_12	35	PURSUIT	0	0.99996	0.00001	0.00003	0.00004	PURSUIT
na030100_13	36	PURSUIT	0	0.99996	0.00002	0.00004	0.00004	PURSUIT
na030100_15	38	PURSUIT	0	0.99994	0.00003	0.00003	0.00005	PURSUIT
na030100_16	39	PURSUIT	0	0.99995	0.00002	0.00003	0.00004	PURSUIT
na030100_18	41	PURSUIT	0	0.99995	0.00002	0.00004	0.00004	PURSUIT
na030100_20	43	PURSUIT	0	0.9999	0.00005	0.00003	0.00008	PURSUIT
na030100_21	44	PURSUIT	0	0.99995	0.00004	0.00004	0.00003	PURSUIT
na030100_22	45	PURSUIT	0	0.99995	0.00002	0.00003	0.00004	PURSUIT
na030600_12	59	Sethoxydim	0.00005	0.00003	0.99701	0	0.00003	Sethoxydim
na030600_14	61	Glyphosate	0.00006	0.00004	0.00002	0.99994	0	Glyphosate
na030600_17	64	Glyphosate	0.00005	0.00005	0.00003	0.99993	0	Glyphosate
na030600_18	65	Foul	0	0.00007	0.00003	0	0.00001	Foul
na030600_19	66	Diuron	0.00034	0.00003	0.00002	0	0.99992	Diuron
na030600_22	69	Diuron	0	0.00005	0.00003	0	0.99989	Diuron
na030600_23	70	Diuron	0.00065	0.00014	0.00043	0	0.92715	Diuron
na030600_24	71	Diuron	0.00002	0.00005	0.00003	0	0.99993	Diuron
Training Set								
na022400_02	2	Control	0.99996	0	0.00001	0.00003	0.00002	Control
na022400_05	5	Control	0.99998	0	0	0.00002	0.00003	Control
na030100_06	29	Control	0.99998	0	0	0.00002	0.00003	Control
na030100_09	32	Control	0.99998	0	0	0.00001	0.00004	Control
na030100_11	34	PURSUIT	0	0.99996	0.00001	0.00003	0.00004	PURSUIT
na030100_14	37	PURSUIT	0	0.99996	0.00001	0.00004	0.00002	PURSUIT
na030100_17	40	PURSUIT	0	0.99995	0.00002	0.00003	0.00004	PURSUIT
na030100_19	42	PURSUIT	0	0.99995	0.00001	0.00004	0.00002	PURSUIT
na030600_04	51	Control	0.99993	0	0.00002	0.00002	0.00001	Control
na030600_08	55	Sethoxydim	0.00002	0.00001	0.99996	0.00005	0.00001	Sethoxydim
na030600_10	57	Sethoxydim	0.00002	0.00002	0.99993	0.00002	0.00001	Sethoxydim
na030600_13	60	Foul	0	0.00004	0.00002	0	0	Foul
na030600_15	62	Glyphosate	0.00007	0.00004	0.00001	0.99992	0	Glyphosate
na030600_16	63	Glyphosate	0.00006	0.00004	0.00003	0.99994	0	Glyphosate
na030600_20	67	Diuron	0.00004	0.00004	0.00002	0	0.99993	Diuron
na030600_21	68	Diuron	0.00007	0.00004	0.00002	0	0.99994	Diuron

Figure 3b

SNNS result file	V1.4-3D
Training file	na022400
Test file	na040400
No. of patterns:	24
No. of input units:	1080
No. of output units:	6
Startpattern:	1
Endpattern:	24
Teaching output included	
Treatment:	1.1 Control
Target:	1 0 0 0 0 0
Output:	0.99954 0.00045 0.00001 0.00001 0.00001 0.00001
Treatment:	2.1 Control
Target:	1 0 0 0 0 0
Output:	0.99936 0.00065 0.00001 0.00001 0.00001 0.00001
Treatment:	3.1 Control
Target:	1 0 0 0 0 0
Output:	0.99951 0.00047 0.00001 0.00001 0.00001 0.00001
Treatment:	4.1 Control
Target:	1 0 0 0 0 0
Output:	0.99963 0.00037 0.00001 0.00001 0.00001 0.00001
Treatment:	5.1 Chlorsulfuron
Target:	0 0 0 0 0 0
Output:	0.00159 0.99843 0 0.00001 0 0
Treatment:	6.1 Chlorsulfuron
Target:	0 0 0 0 0 0
Output:	0.00806 0.99165 0 0 0 0
Treatment:	7.1 Chlorsulfuron
Target:	0 0 0 0 0 0
Output:	0.00334 0.99669 0 0 0 0
Treatment:	8.1 Chlorsulfuron
Target:	0 0 0 0 0 0
Output:	0.00014 0.99985 0 0.00001 0 0
Treatment:	9.1 Chlorsulfuron
Target:	0 0 0 0 0 0
Output:	0.00667 0.99376 0 0 0 0
Treatment:	10.1 Imazamethabenz
Target:	0 0 0 0 0 0
Output:	0.00044 0.99955 0 0 0 0
Treatment:	11.1 Imazamethabenz
Target:	0 0 0 0 0 0
Output:	0.00013 0.99987 0 0 0 0
Treatment:	12.1 Imazamethabenz
Target:	0 0 0 0 0 0
Output:	0.00208 0.99798 0 0.00001 0.00001 0

Figure 4a

SNNS result file	V1.4-3D
Training file	na022400
Test file	na040400
No. of patterns:	24
No. of input units:	1080
No. of output units:	6
Startpattern:	1
Endpattern:	24
Teaching output included	
Treatment:	13.1 Imazamethabenz
Target:	0 0 0 0 0 0
Output:	0.00223 0.99755 0 0 0 0
Treatment:	14.1 Imazamethabenz
Target:	0 0 0 0 0 0
Output:	0.06789 0.93484 0 0 0 0
Treatment:	15.1 Sulfumeturon
Target:	0 0 0 0 0 0
Output:	0.00046 0.99955 0 0 0 0
Treatment:	16.1 Sulfumeturon
Target:	0 0 0 0 0 0
Output:	0.00102 0.999 0 0.00001 0 0
Treatment:	17.1 Sulfumeturon
Target:	0 0 0 0 0 0
Output:	0.00194 0.99813 0 0.00001 0 0
Treatment:	18.1 Sulfumeturon
Target:	0 0 0 0 0 0
Output:	0.00013 0.99987 0 0 0 0
Treatment:	19.1 Sulfumeturon
Target:	0 0 0 0 0 0
Output:	0.00014 0.99985 0 0 0 0
Treatment:	20.1 Imazapyr
Target:	0 0 0 0 0 0
Output:	0.0018 0.998 0 0 0 0
Treatment:	21.1 Imazapyr
Target:	0 0 0 0 0 0
Output:	0.00031 0.99968 0 0.00001 0 0
Treatment:	22.1 Imazapyr
Target:	0 0 0 0 0 0
Output:	0.00175 0.99791 0 0 0 0
Treatment:	23.1 Imazapyr
Target:	0 0 0 0 0 0
Output:	0.00018 0.9998 0 0 0 0
Treatment:	24.1 Imazapyr
Target:	0 0 0 0 0 0
Output:	0.06579 0.93074 0 0 0 0

Figure 4b

Class	Control	AHAS	ACCase	EPSPS	PS II	Necrotic*	HPPD	PROTOX	Carotenoid	PSI	Microtubule	PDS	Uncoupler	Auxin-like	Auxin Transp	DHP	Acetamide	PSII_c1	PSII_c2	PSII_c3	Glutamine	Mitosis	Unknown	No Class
	54	30	6	4	6	2	10	12	9	7	5	9	11	8	1	1	6	9	1	10	12	22	8	4
Control																								
AHAS																								
ACCase																								
EPSPS																								
PS II																								
Necrotic*																								
HPPD																								
PROTOX																								
Carotenoid																								
PSI																								
Microtubule																								
PDS																								
Uncoupler																								
Auxin-like																								
Auxin Transp																								
DHP																								
Acetamide																								
PSII_c1																								
PSII_c2																								
PSII_c3																								
Glutamine																								
Mitosis																								
Unknown																								
No Class																								

Rows: Teaching Input; Columns: Classification for Nineteen MOAs
 (23 Classes, Including "Control" = Untreated Plants, 1 "Spore" Class, and "Unknown" Added by SNNS)

Figure 5

CLASS	Control	AHAS	ACCAse	EPPSPs	HPPD	HPDD	Neurotoxic*	Not Used	PDS	PSI	Carotenoid	PROTOX	HPPD	Auxin-like	Uncoupler	Microtubule	PDS	Uncoupler	Auxin-like	Auxin Transp	DHP	Acetamide	Mitosis	Glutamine	Spare	Unknown	# Plants	
Control	27	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	31	
AHAS	0	17	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	17
ACCase	0	0	3	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	3
EPPSPs	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	2
Not Used	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Necrotic*	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1
HPPD	0	0	0	0	3	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	6
PROTOX	0	0	0	0	0	6	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	6
Carotenoid	0	0	0	0	0	0	6	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	6
PSI	0	0	0	0	0	0	4	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	6
Microtubule	0	0	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	6
PDS	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	3
Uncoupler	0	0	0	0	0	0	0	0	0	4	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	6
Auxin-like	0	0	0	0	0	0	0	0	0	0	0	0	0	4	0	0	0	0	0	0	0	0	0	0	0	0	0	6
Auxin Transp	0	0	0	0	0	0	0	0	0	0	0	0	0	0	3	1	0	0	0	0	0	0	0	0	0	0	0	6
DHP	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	5
Acetamide	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	1	3
PSII_c1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	3	1	0	0	0	0	0	0	0	2	6
PSII_c2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	8	0	0	0	0	0	0	0	1	10
PSII_c3	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	6	0	0	0	0	0	0	0	6
Glutamine	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	11	0	0	0	0	0	0	0	12	
Mitosis	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	3	6
Spare	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
NoClass	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0

Rows: Teaching Input; Columns: Classifications for Nineteen MOAs
 (23 Classes, Including "Control" = Untreated Plants, 1 "Spare" Class, and "Unknown" Added by SNNS)

Figure 6

MOA	Control	AHAS	ACCase	EPSPS	PS II	Necrotic*	HPPD	PROTOX	Carotenoid	PS I	Microtubule	PDS	Cytochrome c	PS II	EPSPS	ACCase	AHAS	Control	Unknown
Control	92	100	100	100	100	100	83	100	100	75	58	83	75	92	67	100	75	17	5.8
AHAS																			
ACCase																			
EPSPS																			
PS II																			
Necrotic*																			
HPPD																			
PROTOX																			
Carotenoid																			
PS I																			
Microtubule																			
PDS																			
Uncoupler																			
Auxin-like																			
Auxin Transp																			
DHP																			
Acetamide																			
PSII_c1																			
PSII_c2																			
PSII_c3																			
Glutamine																			
Mitosis																			
Spare																			
Unknown																			

Rows: Teaching Input, Columns: Classification as Percentage of Total Plants Tested for Nineteen MOAs
 (23 Classes, Including "Control" = Untreated Plants, 1 "Spare" Class, and "Unknown" Added by SNNS)

*Necrotic = dead or decaying plant

Figure 7

CLASS	Class	Control	AHAS	ACCase	EPSPS	Not Used	Necrotic*	HPPD	PROTOX	PDS	PSI	Microtubule	Auxin Transp	Auxin-like	Acetamide	DHP	PSII_c1	PSII_c2	PSII_c3	Glutamine	Mitosis	Spare	Unknown	
Control	0	87																					6	
AHAS	1	100																						
ACCase	2	100																						
EPSPS	3	100																						
Not Used	4																							
Necrotic*	5						100																	
HPPD	6							50																50
PROTOX	7								100															
Carotenoid	8									100														
PSI	9										67													
Microtubule	10										33													33
PDS	11											33												67
Uncoupler	12												67											67
Auxin-like	13													67										17
Auxin	14														50									17
Transp																17								33
DHP	15																							83
Acetamide	16																67							33
PSII_c1	17																	50	17					33
PSII_c2	18																		10	8				10
PSII_c3	19																			100				
Glutamine	20																				92			50
Mitosis	21																							50

Rows: Teaching Input, Columns: Classification as Percentage of Total Plants Tested for Nineteen MOAs
 (23 Classes, Including "Control" = Untreated Plants, 1 "Spare" Class, and "Unknown" Added by SNNS)

Figure 8